IN THE CLAIMS

The following is a listing of the claims in the application.

LISTING OF CLAIMS

- 1. (Cancelled).
- (Currently Amended): A process for preparing a humanized antibody consisting of the steps of:
- (a) <u>first</u> replacing each amino acid residue in the complementarily determining region (CDR) of murine monoclonal antibody heavy chain and light chain variable regions with alanine to produce transformants, selecting a transformant that has a lower affinity to the human antigen (K_D) than the original murine antibody, and determining the replaced amino acid residue of said transformant as a specificity determining residue (SDR); and
- (b) <u>subsequently</u> grafting said SDR to at least one of the corresponding amino acid residues into human antibody variable regions.
- 3. (Previously amended): The process of claim 2, wherein the CDR is selected from the group consisting of HCDR1(aa 31-35), HCDR2(aa 50-65) and HCDR3(aa 95-102) of the heavy chain (SEQ ID NO: 2); and LCDR1(aa 24-34), LCDR2(aa 50-56) and LCDR3(aa 89-97) of the light chain (SEQ ID NO: 4) of the murine monoclonal antibody variable regions that bind hepatitis B virus pre-S1

5

502667024.1

antigen, selecting a transformant that has an affinity to antigen which is more than 3 times lower than the original murine antibody when replaced with alanine, determining the replaced amino acid residue of said transformant as an SDR, and grafting said SDR to the corresponding amino acid sequence in human antibody heavy chain and light chain.

- 4. (Original): The process of claim 3, which is characterized in that the at least one of Trp33, Met34, and Asn35 of HCDRI; Arg50, Tyr52, and Pro52a of HCDR2; and G1u95, Tyr96, and G1u98 of HCDR3 of the murine monoclonal antibody KR127 heavy chain, is grafted to the corresponding amino acid sequences in human antibody heavy chain.
- (Original): The process of claim 4, which is characterized in that the at least one of the following grafting steps is carried out:

the amino acid residue at position 32 in HCDR1 of human antibody with alanine:

the amino acid residue at position 97 in HCDR3 of human antibody with arginine or alanine;

the amino acid residue at position 98 in HCDR3 of human 5 antibody with valine: and

 (d) the amino acid residue at position 102 in HCDR3 of human antibody with arginine or alanine.

502667024.1 6

- 6. (Original): The process of claim 5, which is characterized in that the at least one of Trp33 and Asn35 of HCDRI; Arg50 and Tyr52 of HCDR2; and Arg95 and Tyr96 of HCDR3 of the murine monoclonal antibody KR1 27 heavy chain, is grafted into the human antibody heavy chain DP7-JH4.
- 7. (Previously amended): The process of claim 6, which is characterized in that the amino acid residues of the A1a71 and Lys73 in Framework region 3 of the murine monoclonal antibody KR127 heavy chain variable region, are further grafted into the human antibody heavy chain DP7-JH4.
- 8. (Original): The process of claim 3, which is characterized in that the at least one of the Leu27b, Tyr27d, Ser27e, Asn28, Lys30, Tyr32 and Asn34 of LCDR1; Leu50 and Asp55 of LCDR2; and Va189, Gln90, G1y91, Thr92, His93, Phe94, Pro95, and G1n96 of LCDR3 of the murine monoclonal antibody KR1 27 light chain, is grafted into the human antibody light chain.
- 9. (Currently Amended): The process of claim 8, which is characterized in that the Tyr27d, Asn28, Asn34 of LCDR1; Leu50 and Asp55 of LCDR2; and Va189, Gly91, Thr92, His93, Phe94, Pro95, and Gln96 of LCDR3 of the murine monoclonal antibody KR127 light chain, is grafted into the human antibody light chain DPH12_JK4 DPK12_JK4.

7

502667024.1

- 10. (Original): The process of claim 8, which is characterized in that the Leu36 and Arg46 in Framework region 2 of the murine monoclonal antibody KR127 light chain variable region, are further grafted into the human antibody light chain DPH12-JK4 DPK12-JK4.
- 11. (Withdrawn): A humanized antibody prepared by the process of any one of claims 1 to 10, which has an affinity to antigen of higher than 8.2 x 1 0⁹ M and suppresses HAMA (human anti-mouse antibody) response to a greater extent than an antibody prepared according to CDR-grafting method.
- 12. (Withdrawn): The humanized antibody of claim 11, which has the amino acid sequence of SEQ ID NO: 2 for the heavy chain variable region of HBV pre-S1 antigen.
- 13. (Withdrawn): The humanized antibody of claim 11, which has the amino acid sequence of SEQ ID NO: 4 for the light chain variable region of HBV pre-S1 antigen.
- 14. (Withdrawn): A humanized antibody which is produced by CHO/HuKR127 (Accession No.: KCTC 10199BP).
- 15. (Withdrawn): A DNA encoding the humanized antibody heavy chain containing the amino acid sequence of SEQ ID NO: 2 for the heavy chain variable region of HBV pre-S1 antigen.

8

502667024.1

- 16. (Withdrawn): The DNA of claim 15, wherein the variable region has the nucleotide sequence of SEQ ID NO: 1.
- 17. (Withdrawn): A DNA encoding the humanized antibody light chain containing the amino acid sequence of SEQ ID NO: 4 for the light chain variable region of HBV pre-S1 antigen.
- 18. (Withdrawn): The DNA of claim 17, wherein the variable region has the nucleotide sequence of SEQ ID NO: 3.
- 19. (Withdrawn): An expression vector pHuKR127HC comprising the DNA of claim 16 for expressing the humanized antibody heavy chain for HBV pre-S1 antigen.
- 20. (Withdrawn): An expression vector pHuKR127KC comprising the DNA of claim 18 for expressing the humanized antibody light chain for HBV pre- S1 antigen.
- 21. (Withdrawn): An expression vector pdCMV-dhfrC-HuKR127 comprising the nucleotide sequences of SEQ ID NO:1 and SEQ ID NO:3 for expressing the humanized antibody light and heavy chains for HBV pre-S1antigen.
- (Withdrawn): An E. coli DH5a/pdCMV-dhfrC-HuKR127 (Accession No.:
 KCTC 10198BP) transformed with the expression vector of claim 21.

502667024.1 9

- 23. (Withdrawn): CHO cell line CHOIHuKR127 (Accession No.: KCTC 1010199BP) producing the humanized antibody of claim 11.
- 24. (Withdrawn): A composition for preventing or treating HBV infection comprising the humanized antibody of claim 11.

502667024.1 10